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SEQUENCE LISTING

<110> ASANO Shinichiro et al.

<120> Protein Having Insecticidal Activity, DNA Coding Said Protein, Pest Control Agent and Pest Control Method

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Ile Thr Gln Ile Pro Val Val Lys Ala His Thr Leu Val Ser Gly Thr
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995 1000 1005

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caa acg acc aca tta caa aac atg aac tat aaa gat tat ctg aga atg 144
Gln Thr Thr Thr Leu Gln Asn Met Asn Tyr Lys Asp Tyr Leu Arg Met
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Ser Glu Gly Glu Asn Pro Glu Leu Phe Gly Asn Pro Glu Thr Phe Ile
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Ser Ser Ser Thr Val Gln Thr Gly Ile Gly Ile Val Gly Gln Val Leu
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Gly Ala Leu Gly Val Pro Phe Ala Gly Gln Ile Ala Ser Phe Tyr Ser
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Phe Ile Val Gly Gln Leu Trp Pro Ser Ser Thr Val Ser Val Trp Glu
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Met Ile Met Lys Gln Val Glu Asp Leu Ile Asp Gln Lys Ile Thr Asp
115 120 125

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Ser Val Arg Lys Thr Ala Leu Ala Gly Leu Gln Gly Leu Gly Asp Gly
130 135 140

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Asp Thr Arg Ala Arg Ser Val Val Val Thr Gln Tyr Ile Ala Leu Glu
165 170 175

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Leu Asp Phe Val Ala Lys Ile Pro Ser Phe Ala Ile Ser Gly Gln Glu
180 185 190

gta cca tta tta tca gtg tat gca caa gca gcg aat tta cat ttg cta 624
Val Pro Leu Leu Ser Val Tyr Ala Gln Ala Ala Asn Leu His Leu Leu
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Tyr Ser Asp Tyr Cys Val Lys Trp Tyr Asn Thr Gly Leu Asp Lys Leu
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aaa ggt acg aat gct gca agt tgg ctg aag tat cac caa ttc cga aga   816
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Glu Met Thr Leu Leu Val Leu Asp Leu Val Ala Leu Phe Pro Asn Tyr
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gac aca cgt acg tat cca atc gaa aca acg gcc caa ctt aca cgg gaa   912
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tgt agg cgt tgg tca ctt aac agt gat att tct ttt tca gaa gtc gaa   1008
Cys Arg Arg Trp Ser Leu Asn Ser Asp Ile Ser Phe Ser Glu Val Glu
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agc gct gta att cgt tca cca cac cta ttt gat ata ctc agt gaa ata   1056
Ser Ala Val Ile Arg Ser Pro His Leu Phe Asp Ile Leu Ser Glu Ile
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tca tca gca tta gaa cgt aat tac ggt acg att act tct aac aaa atc   1200
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aag tat tat gat tta gca aat aag gat atc ttt cag gtt cga tca tta   1248
Lys Tyr Tyr Asp Leu Ala Asn Lys Asp Ile Phe Gln Val Arg Ser Leu
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ggg gcg gat tta gct aat tac tac gca cag gta tat gga gtt ccg tac   1296
Gly Ala Asp Leu Ala Asn Tyr Tyr Ala Gln Val Tyr Gly Val Pro Tyr
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gct agt ttt aca ctg ctt gac aag aat aca gga tca gga tca gtt gga   1344
Ala Ser Phe Thr Leu Leu Asp Lys Asn Thr Gly Ser Gly Ser Val Gly
        435                 440                 445

ggt ttt acg tac tca aaa cca cat aca act atg caa gta tgt aca caa   1392
Gly Phe Thr Tyr Ser Lys Pro His Thr Thr Met Gln Val Cys Thr Gln
    450                 455                 460

aat tac aat acg att gat gaa atc cct cca gag aat gag cca ctt agt   1440
Asn Tyr Asn Thr Ile Asp Glu Ile Pro Pro Glu Asn Glu Pro Leu Ser
465                 470                 475                 480

aga ggg tat agc cat aga tta tct cat atc acc tct tat tct ttt tct   1488
Arg Gly Tyr Ser His Arg Leu Ser His Ile Thr Ser Tyr Ser Phe Ser
                485                 490                 495

aag aat gct agt agt cct gct aga tat ggc aat ctc cct gta ttt gct   1536
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att act cag ata cca gtt gta aag gca cat act tta gtt tca ggt act   1632
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    530                 535                 540

act gtt att aaa ggt cct gga ttt aca gga ggc aat atc ctt aaa aga   1680
Thr Val Ile Lys Gly Pro Gly Phe Thr Gly Gly Asn Ile Leu Lys Arg
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aca agt agt ggt ccg tta gct tat act agt gtc tct gta aaa tca cca   1728
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tta tca caa aga tat cgt gca aga ata cgt tat gct tct act act aac   1776
Leu Ser Gln Arg Tyr Arg Ala Arg Ile Arg Tyr Ala Ser Thr Thr Asn
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tta cga ctt ttt gta aca att tct gga act cgc att tac tct ata aat   1824
Leu Arg Leu Phe Val Thr Ile Ser Gly Thr Arg Ile Tyr Ser Ile Asn
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gtt aat aaa acc atg aat aaa ggg gat gat tta aca ttt aat aca ttt   1872
Val Asn Lys Thr Met Asn Lys Gly Asp Asp Leu Thr Phe Asn Thr Phe
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gac tta gca act att ggt act gct ttc aca ttt tca aat tac tcg gat   1920
Asp Leu Ala Thr Ile Gly Thr Ala Phe Thr Phe Ser Asn Tyr Ser Asp
625                 630                 635                 640

agc tta acg gta ggt gca gat tct ttt gct tca gga gga gaa gtt tat   1968
Ser Leu Thr Val Gly Ala Asp Ser Phe Ala Ser Gly Gly Glu Val Tyr
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gta gat aag ttc gaa ctt att ccg gta aat gca aca ttt gaa gca gaa   2016
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Lys Lys Asp Ala Leu Gln Thr Ser Val Thr Asp Tyr Gln Val Asn Gln
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gcg gca aac tta gta gaa tgc cta tcc gat gag tta tac cca aat gaa   2160
Ala Ala Asn Leu Val Glu Cys Leu Ser Asp Glu Leu Tyr Pro Asn Glu
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aaa cga atg tta tgg gat gca gtg aaa gag gcg aaa cga ctt gtt cag   2208
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gca cgt aac tta ctc caa gat aca ggc ttt aat agg att aat gga gaa   2256
Ala Arg Asn Leu Leu Gln Asp Thr Gly Phe Asn Arg Ile Asn Gly Glu
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aac gga tgg acg gga agt acg gga atc gag gtt gcg gaa gga gat gtt   2304
Asn Gly Trp Thr Gly Ser Thr Gly Ile Glu Val Ala Glu Gly Asp Val
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ctg ttt aaa gat cgt tcg ctt cgt ttg aca agt gcg aga gag att gat   2352
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785                 790                 795                 800

tta aaa cca tat aca aga tat aaa cta aaa ggt ttt ata gga agt agt   2448
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caa gat tta gag att aaa tta ata cgt cat cgg gca aat caa atc gtc 2496
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835 840 845

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tcc att cct tac gta tat aat gat gcg tta cca gaa atc cct gga atg   3024
Ser Ile Pro Tyr Val Tyr Asn Asp Ala Leu Pro Glu Ile Pro Gly Met
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aac tat acg agt ttt aca gag tta aca aat aga ctc caa caa gca tgg   3072
Asn Tyr Thr Ser Phe Thr Glu Leu Thr Asn Arg Leu Gln Gln Ala Trp
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aat ttg tat gat ctt cga aat gct ata cca aat gga gat ttt cga aat   3120
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agc gat aca tct gtc ctt gtc att cca aac tgg aat tct caa gtg tca   3216
Ser Asp Thr Ser Val Leu Val Ile Pro Asn Trp Asn Ser Gln Val Ser
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caa caa ttt aca gtt caa ccg aat tat aga tat gtg tta cgt gtc aca   3264
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1140 1145 1150

ggt gta ttc aac ata gaa agt gta gaa ctc gtg tta gaa gaa gag taa 3504
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